

Figure 1

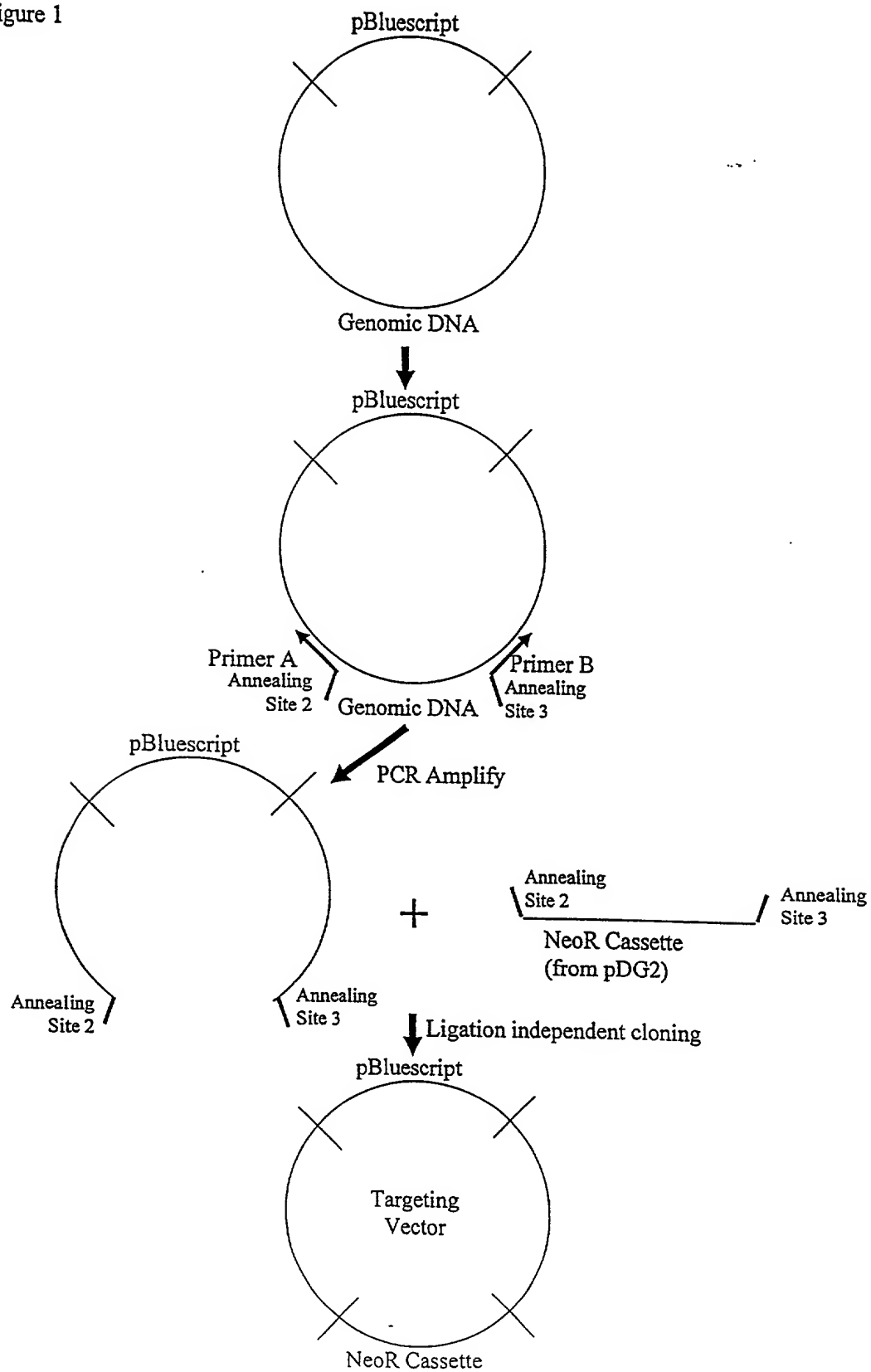
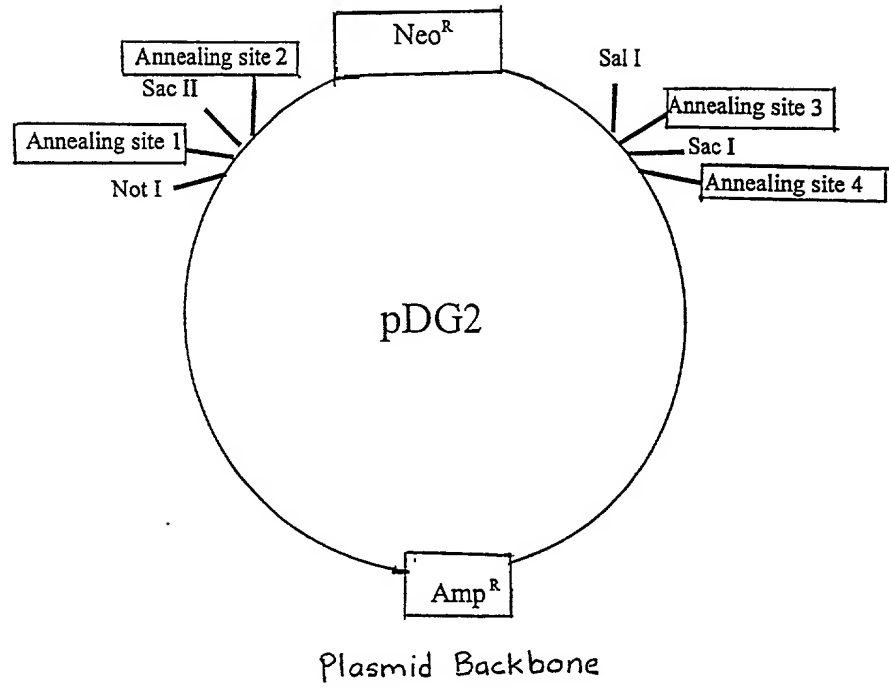


Figure 2A



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Fig 2B

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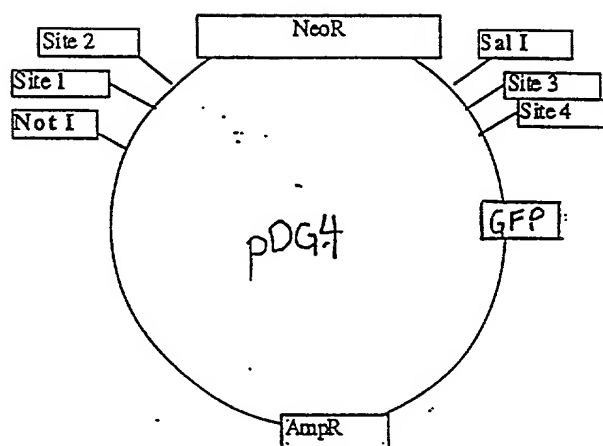


Fig 3A

REPORT

Fig 3B

20220123/2001

Annealing site	Sequence	Sequence after digestion
1	5' tgtgctcctctttggcttgcttcaa... 3' 3' acacgaggagaaaaccgaacgaagggt... 5'	5' tgtgctcctctttggcttgcttcaa... 3' 3' tt... 5'
2	5' ctggttcttgtctggttggcccaa... 3' 3' gaccaagaacagaccgaacgggtt... 5'	5' ctggttcttgtctggttggcccaa... 3' 3' tt... 5'
3	5' ggtcctcgctctgtgtccgttgaa... 3' 3' ccaggagcgagacacaggcaactt... 5'	5' ggtcctcgctctgtgtccgttgaa... 3' 3' tt... 5'
4	5' tttgcgtgtcctgtgtcgtcgaa... 3' 3' aaacgcacaggacacagcagctt... 5'	5' tttgcgtgtcctgtgtcgtcgaa... 3' 3' tt... 5'

Fig 4

Fig 5

Annealing site	Sequence	Sequence after digestion
1	5' AATgtgctcctcttttggttgettCOGC 3' 3' Ttacacgaggagaaaccgaacgaagg 5'	5' AA 3' 3' Ttacacgaggagaaaccgaacgaagg 5'
2	5' AActggttcttgtctggcttggcCOGC 3' 3' Ttgaccaagaacagaccgaaccggg 5'	5' AA 3' 3' Ttgaccaagaacagaccgaaccggg 5'
3	5' AAggtcctcgctctgtgtcgttGAGCT 3' 3' Ttccaggagcgagacacaggcaac 5'	5' AA 3' 3' Ttccaggagcgagacacaggcaac 5'
4	5' AAtttgcgtgtcctgtgtcgtcGAGCT 3' 3' Ttaaacgcacaggacacagcagc 5'	5' AA 3' 3' Ttaaacgcacaggacacagcagc 5'

2003.03.20 14:23:00

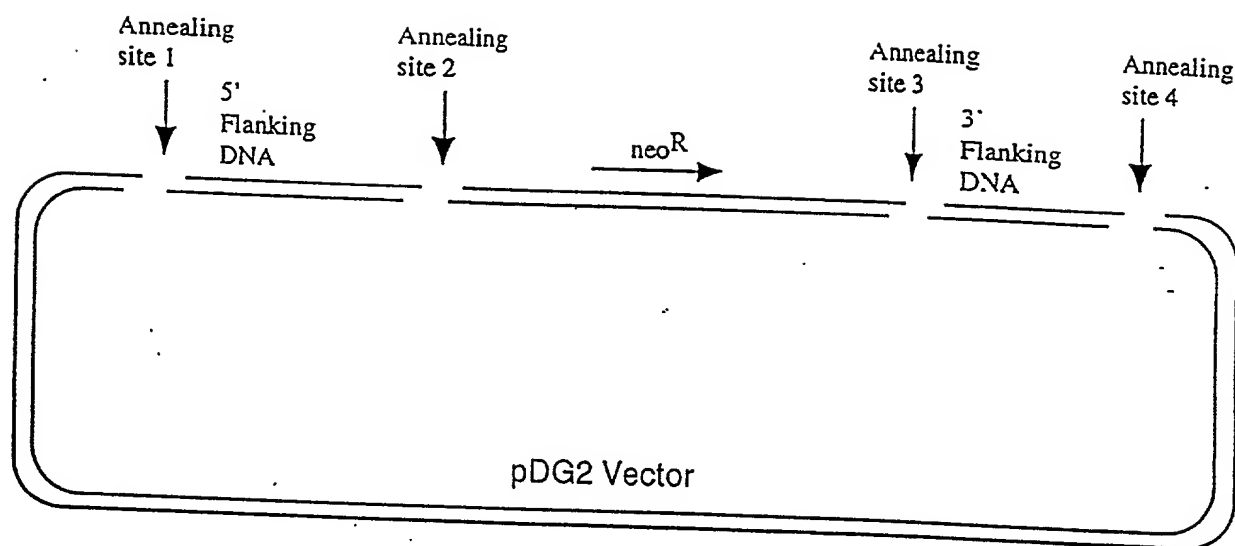


Fig 6

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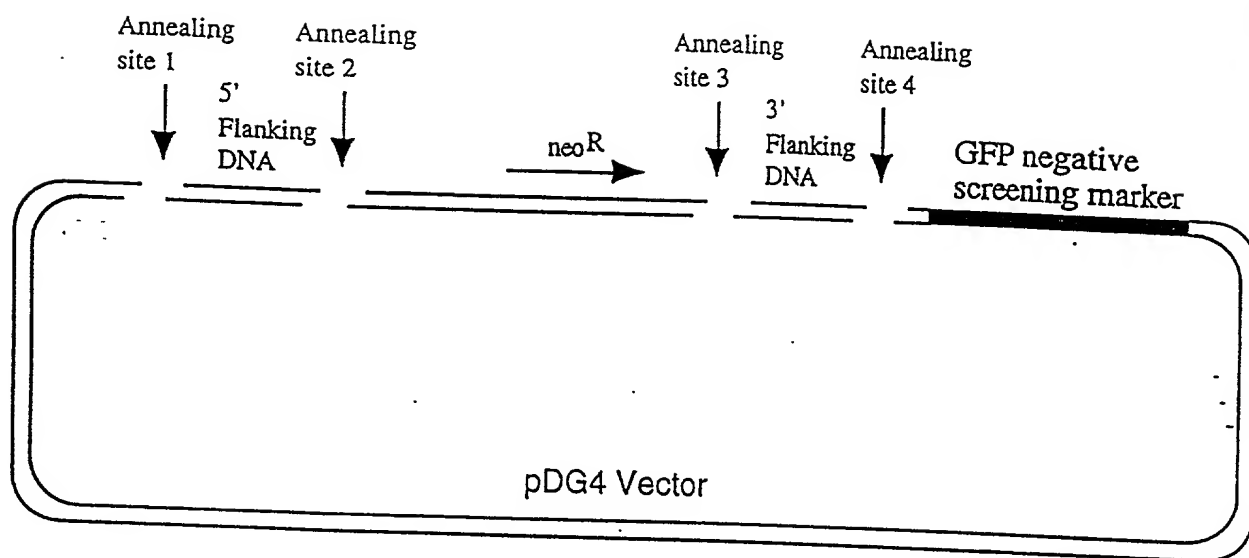


Fig 7

Oligo#	Sequence (5' to 3')
174	ATGACCGCTCAGGAAACCTGTTGCA
180	ATAGGCATAGTAGGCCAGCTTGAGG
454	tgtgctcctctttggcttgcttccAATTAACCCTCACTAAAGGGAACGAAT
463	ctgggttcttgtctggcttggcccaaTGCAACAGGTTTCCTGAGCGGTCAT
464	ggtcctcgctctgtgtccggttgaaCCTCAAGCTGGCCTACTATGCCTAT
42	tttgctgtcctgtgtcgctcgaaCGACTAATACGACTCACTATAGGGCG
151	GCCAATGGACTCTTAGTTTTGGAAC
155	GTTCTGGCAAACAAATTCGGCGCAC
454	tgtgctcctctttggcttgcttccAATTAACCCTCACTAAAGGGAACGAAT
465	ctgggttcttgtctggcttggcccaaGTTCCAAACTAAGAGTCCATTGGC
466	ggtcctcgctctgtgtccggttgaaGTGCGCCGAATTTGTTTGCCAGAAC
1	GAACCTTGGTGTGCCAAGTTACTTC
2	GAACCTTGGCTGAACCCCTTGTCT
41	tgtgctcctctttggcttgcttgaCGACTAATACGACTCACTATAGGGCG
38	ctgggttcttgtctggcttggcccaaGAAGTAAGTTGGCACACCAAGGTC
40	ggtcctcgctctgtgtccggttgaaGAACAAGGGGTTAGCCAAAGTTC
37	tttgctgtcctgtgtcgctcgAATTAACCCTCACTAAAGGGAACGAAT
540	ATGCCGGATCTCTACTACTGGGCC
546	TGTCATAGTAGACAGCGATGGAACG
445	GACAAGAACCAGTTGACGTCAAGCTTCCCGGACGCGTGCTAGCGGCGCGCCG
667	ctgggttcttgtctggcttggcccaaGGCCCAGTAGTAGGAGATCCGGCAT
668	ggtcctcgctctgtgtccggttgaaCGTTCCATCGCTGTCTACTATGACA
907	ctgggttcttgtctggcttggcccaaAAAGCCGACAGCCACGCTCACAAGC
908	ggtcctcgctctgtgtccggttgaaGCCCAATGCCACAGAGACAGAATGT
1157	ctgggttcttgtctggcttggcccaaGTTGGATCCTCTCCAAGGCCCCATCT
1158	ggtcctcgctctgtgtccggttgaaCTCCAGTGCCGAGTGTGTGGGGACAG

Figure 8